<110> Yu et al. <120> Neutrokine-alpha and Neutrokine-alpha Splice Variants <130> PF343P3 <140> 09/507,968 <141> 2000-02-22 <150> 60/122,388 <151> 1999-03-02 <150> 60/124,097 <151> 1999-03-12 <150> 60/126,599 <151> 1999-03-26 <150> 60/127,598 <151> 1999-04-02 <150> 60/130,412 <151> 1999-04-16 <150> 60/130,696 <151> 1999-04-23 <150> 60/131,278 <151> 1999-04-27 <150> 09/255,794 <151> 1999-02-23 <150> 60/131,673 <151> 1999-04-29 <150> 60/136,784 <151> 1999-05-28 <150> 60/142,659 <151> 1999-07-06 <150> 60/145,824 <151> 1999-07-27

<150> 60/168,624

<150> 60/167,239 <151> 1999-11-24

<151> 1999-12-03

<150> 60/171,108 <151> 1999-12-16

<150> 60/171,626

<151> 1999-12-23

<150> 60/176,015

<151> 2000-01-14 <160> 38 <170> PatentIn Ver. 2.1 <210> 1 <211> 1100 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (147)..(1001) aaattcagga taactctcct \gaggggtgag ccaagccctg ccatgtagtg cacgcaggac 60 atcaacaaac acagataaca gaaaatgatc cattccctgt ggtcacttat tctaaaggcc 120 ccaaccttca aagttcaagt agy gat atg gat gac tcc aca gaa agg gag cag Met Asp Asp Ser Thr Glu Arg Glu Gln tea ege ett aet tet tge ett aag aaa aga gaa gaa atg aaa etg aag 221 Ser Arg Leu Thr Ser Cys Leu Lys Lys Arg Glu Glu Met Lys Leu Lys 15 gag tgt gtt tcc atc ctc cca cgg\aag gaa agc ccc tct gtc cga tcc 269 Glu Cys Val Ser Ile Leu Pro Arg Lys Glu Ser Pro Ser Val Arg Ser tee aaa gae gga aag etg etg get gda ace ttg etg etg gea etg etg 317 Ser Lys Asp Gly Lys Leu Leu Ala Ala Thr Leu Leu Leu Ala Leu Leu tet tgc tgc etc acg gtg gtg tet ttc tac cag gtg gcc gcc etg caa 365 Ser Cys Cys Leu Thr Val Val Ser Phe Tyr Gln Val Ala Ala Leu Gln 65 ggg gac ctg gcc agc ctc cgg gca gag ctg cag ggc cac cac gcg gag 413 Gly Asp Leu Ala Ser Leu Arg Ala Glu Leu Gln Gly His His Ala Glu aag ctg cca gca gga gca gga gcc ccc aag g|cc ggc ctg gag gaa gct 461 Lys Leu Pro Ala Gly Ala Gly Ala Pro Lys Ala Gly Leu Glu Glu Ala 90 100 cca gct gtc acc gcg gga ctg aaa atc ttt gaa cca cca gct cca gga 509 Pro Ala Val Thr Ala Gly Leu Lys Ile Phe Glu/Pro Pro Ala Pro Gly 115 120 gaa ggc aac tcc agt cag aac agc aga aat aag dgt gcc gtt cag ggt 557 Glu Gly Asn Ser Ser Gln Asn Ser Arg Asn Lys Arg Ala Val Gln Gly 125 130 cca gaa gaa aca gtc act caa gac tgc ttg caa ctd att gca gac agt 605 Pro Glu Glu Thr Val Thr Gln Asp Cys Leu Gln Leu Ile Ala Asp Ser 140 145

.

| | | | | ١ ١ | | | | | | | | | | | | |
|------------------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-----------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| gaa Glu | aca Thr 155 | Pro | act Thr | ata Ile | Caa Gln | aaa Lys 160 | Gly | tct Ser | tac Tyr | aca Thr | ttt Phe 165 | Val | cca Pro | tgg Trp | ctt Leu | 653 |
| ctc Leu 170 | agc Ser | ttt Phe | aaa Lys | agg Arg | gga Gly 175 | agt Ser | gcc Ala | cta Leu | gaa Glu | gaa Glu 180 | aaa Lys | gag Glu | aat Asn | aaa Lys | ata Ile 185 | 701 |
| ttg Leu | gtc Val | aaa Lys | gaa Glu | act Thr 190 | GJA GGA | tac Tyr | ttt Phe | ttt Phe | ata Ile 195 | tat Tyr | ggt Gly | cag Gln | gtt Val | tta Leu 200 | tat Tyr | 749 |
| act Thr | gat Asp | aag Lys | acc Thr 205 | tac Tyr | gcc Ala | Met | gga Gly | cat His 210 | cta Leu | att Ile | cag Gln | agg Arg | aag Lys 215 | aag Lys | gtc Val | 797 |
| cat His | gtc Val | ttt Phe 220 | gly aaa | gat Asp | gaa Glu | t tig Leu | agt Ser 225 | ctg Leu | gtg Val | act Thr | ttg Leu | ttt Phe 230 | cga Arg | tgt Cys | att Ile | 845 |
| caa Gln | aat Asn 235 | atg Met | cct Pro | gaa Glu | aca Thr | cta Leu 240 | ccc | aat Asn | aat Asn | tcc Ser | tgc Cys 245 | tat Tyr | tca Ser | gct Ala | ggc Gly | 893 |
| att Ile 250 | gca Ala | aaa Lys | ctg Leu | gaa Glu | gaa Glu 255 | gga Gly | gat Asp | gaa Glu | ctc Leu | caa Gln 260 | ctt Leu | gca Ala | ata Ile | cca Pro | aga Arg 265 | 941 |
| gaa Glu | aat Asn | gca Ala | caa Gln | ata Ile 270 | tca Ser | ctg Leu | gat Asp | gga Gly | gat Asp 275 | gtc Val | aca Thr | ttt Phe | ttt Phe | ggt Gly | gca Ala 280 | 989 |
| ttg Leu | | | | tgad | ctac | ett a | cacc | atbt | c tg | rtago | tatt | ttc | ctcc | ctt | 285 | 1041 |
| tctc | tgta | cc t | ctaa | gaag | ja aa | ıgaat | ctaa | ctg | aaaa | itac | caaa | aaaa | aa a | aaaa | .aaaa | 1100 |
| <210 <211 <212 <213 | > 28 > PR | T | apie | ns | | | | | | | | | | 1 | | |
| <400: Met / | | Asp | Ser | Thr 5 | Glu | Arg | Glu | Gln | Ser 10 | Arg : | Leu | Thr | Ser | Cys 15 | Leu | |
| Lys 1 | Lys | Arg | Glu 20 | Glu | Met | Lys | Leu | Lys 25 | Glu | ¢γs . | Val | Ser | Ile 30 | Leu | Pro | |
| Arg] | Ĺуs | Glu 35 | Ser : | Pro | Ser | Val | Arg 40 | Ser | Ser | Lys i | Asp | Gly : | Lys : | Leu | Leu | |
| Ala A | Ala 50 | Thr | Leu : | Leu | Leu | Ala 55 | Leu | Leu : | Ser | Cys | Cys : | Leu ' | Thr ' | Val ' | Val | |
| Ser 1 65 | Phe ' | Tyr | Gln ' | Val . | Ala 70 | | Leu (| Gln (| Gly . | Asp } | Leu I | Ala : | Ser 1 | Leu . | Arg 80 | |

Ala Glu Leu Gln Gly His His Ala Glu Lys Leu Pro Ala Gly Ala Gly 90 Ala Pro Lys Ala Gly Leu Glu Glu Ala Pro Ala Val Thr Ala Gly Leu 100 105 Lys Ile Phe Glu Pro Pto Ala Pro Gly Glu Gly Asn Ser Ser Gln Asn Ser Arg Asn Lys Arg Ala\ Val Gln Gly Pro Glu Glu Thr Val Thr Gln 135 Asp Cys Leu Gln Leu Ile Ala Asp Ser Glu Thr Pro Thr Ile Gln Lys 150 Gly Ser Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys Arg Gly Ser 165 170 Ala Leu Glu Glu Lys Glu Asn Lys Ile Leu Val Lys Glu Thr Gly Tyr 185 Phe Phe Ile Tyr Gly Gln Val Leu Tyr Thr Asp Lys Thr Tyr Ala Met Gly His Leu Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu Ser Leu Val Thr Leu Phe Arg Cys tle Gln Asn Met Pro Glu Thr Leu 225 230 Pro Asn Asn Ser Cys Tyr Ser Ala Gly Ile Ala Lys Leu Glu Gly Asp Glu Leu Gln Leu Ala Ile Pro Arg\Glu Asn Ala Gln Ile Ser Leu Asp Gly Asp Val Thr Phe Phe Gly Ala Leu Lys Leu Leu 275 280

<210> 3 <211> 233 <212> PRT <213> Homo sapiens

<400> 3 Met Ser Thr Glu Ser Met Ile Arg Asp Val Glu Leu Ala Glu Glu Ala

Leu Pro Lys Lys Thr Gly Gly Pro Gln Gly Set Arg Arg Cys Leu Phe

Leu Ser Leu Phe Ser Phe Leu Ile Val Ala Gly Ala Thr Thr Leu Phe

Cys Leu Leu His Phe Gly Val Ile Gly Pro Gln Arg Glu Glu Phe Pro

Arg Asp Leu Ser Leu Ile Ser Pro Leu Ala Gln Ala Val Arg Ser Ser

75

| Ser | Arg | Thr | Pro | Ser | Asp | Lys | Pro | Val | Ala | His | Val | Val | Ala | Asn | Pro |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | | | 85 | \ | | | | 90 | | | | | 95 | |

Gln Ala Glu Gly Gln Leu Gln Trp Leu Asn Arg Arg Ala Asn Ala Leu
100 105 110

Leu Ala Asn Gly Val Glu Leu Arg Asp Asn Gln Leu Val Val Pro Ser 115 120 125

Glu Gly Leu Tyr Leu Ile Tyr Ser Gln Val Leu Phe Lys Gly Gln Gly - 130 140

Cys Pro Ser Thr His Val Leu Thr His Thr Ile Ser Arg Ile Ala 145 150 155 160

Val Ser Tyr Gln Thr Lys Val Asn Leu Leu Ser Ala Ile Lys Ser Pro 165 170 175

Cys Gln Arg Glu Thr Pro Glu Gly Ala Glu Ala Lys Pro Trp Tyr Glu
180

Pro Ile Tyr Leu Gly Gly Val Phe Gln Leu Glu Lys Gly Asp Arg Leu
195 200 205

Ser Ala Glu Ile Asn Arg Pro Asp Tyr Leu Asp Phe Ala Glu Ser Gly 210 220

Gln Val Tyr Phe Gly Ile Ile Ala Leu 225 230

<210> 4

nocowe no och

<211> 205

<212> PRT

<213> Homo sapiens

<400> 4

Met Thr Pro Pro Glu Arg Leu Phe Leu Pro Arg Val Arg Gly Thr Thr
1 5 10 15

Leu His Leu Leu Leu Gly Leu Leu Leu Val Leu Leu Pro Gly Ala
20 25 30

Gln Gly Leu Pro Gly Val Gly Leu Thr Pro Ser\Ala Ala Gln Thr Ala
35 40 45

Arg Gln His Pro Lys Met His Leu Ala His Ser Thr Leu Lys Pro Ala

Ala His Leu Ile Gly Asp Pro Ser Lys Gln Asn Ser Leu Leu Trp Arg
65 70 75 80

Ala Asn Thr Asp Arg Ala Phe Leu Gln Asp Gly Phe Ser Leu Ser Asn 85 90 95

Asn Ser Leu Leu Val Pro Thr Ser Gly Ile Tyr Phe Val Tyr Ser Gln
100 105 110

Val Val Phe Ser G∦y Lys Ala Tyr Ser Pro Lys Ala Thr Ser Ser Pro 120 Leu Tyr Leu Ala His Glu Val Gln Leu Phe Ser Ser Gln Tyr Pro Phe 135 His Val Pro Leu Leu Ser Ser Gln Lys Met Val Tyr Pro Gly Leu Gln 1/20 Glu Pro Trp Leu His Ser Met Tyr His Gly Ala Ala Phe Gln Leu Thr 170 Gln Gly Asp Gln Leu Ser Thr His Thr Asp Gly Ile Pro His Leu Val 185 Leu Ser Pro Ser Thr Val Phe Phe Gly Ala Phe Ala Leu 195 200 <210> 5 <211> 244 <212> PRT <213> Homo sapiens <400> 5 Met Gly Ala Leu Gly Leu Glu Gly Arg Gly Gly Arg Leu Gln Gly Arg 5 10 Gly Ser Leu Leu Ala Val Ala Gly Ala Thr Ser Leu Val Thr Leu Leu Leu Ala Val Pro Ile Thr Val Leu Ala Val Leu Ala Leu Val Pro Gln Asp Gln Gly Gly Leu Val Thr Gl\(\frac{1}{4}\) Thr Ala Asp Pro Gly Ala Gln Ala Gln Gln Gly Leu Gly Phe Gln Lys teu Pro Glu Glu Glu Pro Glu 65 Thr Asp Leu Ser Pro Gly Leu Pro Ala Ala His Leu Ile Gly Ala Pro Leu Lys Gly Gln Gly Leu Gly Trp Glu Thr Thr Lys Glu Gln Ala Phe 105 Leu Thr Ser Gly Thr Gln Phe Ser Asp Ala Glu Gly Leu Ala Leu Pro 120 Gln Asp Gly Leu Tyr Tyr Leu Tyr Cys Leu Vall Gly Tyr Arg Gly Arg 135 Ala Pro Pro Gly Gly Gly Asp Pro Gln Gly Arg\Ser Val Thr Leu Arg Ser Ser Leu Tyr Arg Ala Gly Gly Ala Tyr Gly Aro Gly Thr Pro Glu 170 Leu Leu Clu Gly Ala Glu Thr Val Thr Pro Val Leu Asp Pro Ala

185

190

180

Arg Arg Gln Gly Tyr Gly Pro Leu Trp Tyr Thr Ser Val Gly Phe Gly 200 Gly Leu Val Gln Leu Arg Arg Gly Glu Arg Val Tyr Val Asn Ile Ser His Pro Asp Met Val Asp Phe Ala Arg Gly Lys Thr Phe Phe Gly Ala 23b Val Met Val Gly <210> 6 <211> 281 <212> PRT <213> Homo sapiens <220> <223> Description of Combined DNA/RNA Molecule: n equals a, t, g, or c <400> 6 Met Gln Gln Pro Phe Asn Tyr Pro Tyr Pro Gln Ile Tyr Trp Val Asp Ser Ser Ala Ser Ser Pro Trp Ala Pro Pro Gly Thr Val Leu Pro Cys Pro Thr Ser Val Pro Arg Arg Prd Gly Gln Arg Arg Pro Pro Pro Pro Pro Pro Pro Leu Pro Pro Pro Pro Pro Pro Pro Leu Pro Pro Leu Pro Leu Pro Pro Leu Lys Lys Arg Gly Asn His Ser Thr Gly 75 Leu Cys Leu Leu Val Met Phe Phe Met Val Leu Val Ala Leu Val Gly Leu Gly Leu Gly Met Phe Gln Leu Phe His Leu Gln Lys Glu Leu Ala 105 Glu Leu Arg Glu Ser Thr Ser Gln Met His Thr Ala Ser Ser Leu Glu 120 Lys Gln Ile Gly His Pro Ser Pro Pro Glu Lys Lys Glu Leu Arg 130 Lys Val Ala His Leu Thr Gly Lys Ser Ash Ser Arg Ser Met Pro Leu 155 150 Glu Trp Glu Asp Thr Tyr Gly Ile Val Leu\Leu Ser Gly Val Lys Tyr

Lys Lys Gly Gly Leu Val Ile Asn Glu Thr Gly Leu Tyr Phe Val Tyr

```
Ser Lys Val Tyr Phe Arg Gly Gln Ser Cys Asn Asn Leu Pro Leu Ser
His Lys Val Tyr Met Arg Asn Ser Lys Tyr Pro Gln Asp Leu Val Met
                         215
Met Glu Gly Lys Met Net Ser Tyr Cys Thr Thr Gly Gln Met Trp Ala
                                         235
Arg Ser Ser Tyr Leu Gly Ala Val Phe Asn Leu Thr Ser Ala Asp His
                245
Leu Tyr Val Asn Val Sex Glu Leu Ser Leu Val Asn Phe Glu Glu Ser
            260
                                 265
Gln Thr Phe Phe Gly Leu Tyr Lys Leu
        275
                             280
<210> 7
<211> 337
<212> DNA
<213> Homo sapiens
<220>
<223> Description of Combined DNA/RNA Molecule: n equals
      a, t, g, or c
<220>
<221> misc_feature
<222> (3)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (58)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (67) . . (71)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (212)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (255)
<223> n equals a, t, g or c
<220>
<221> misc feature
<222> (297)
<223> n equals a, t, g or c
<220>
<221> misc_feature
```

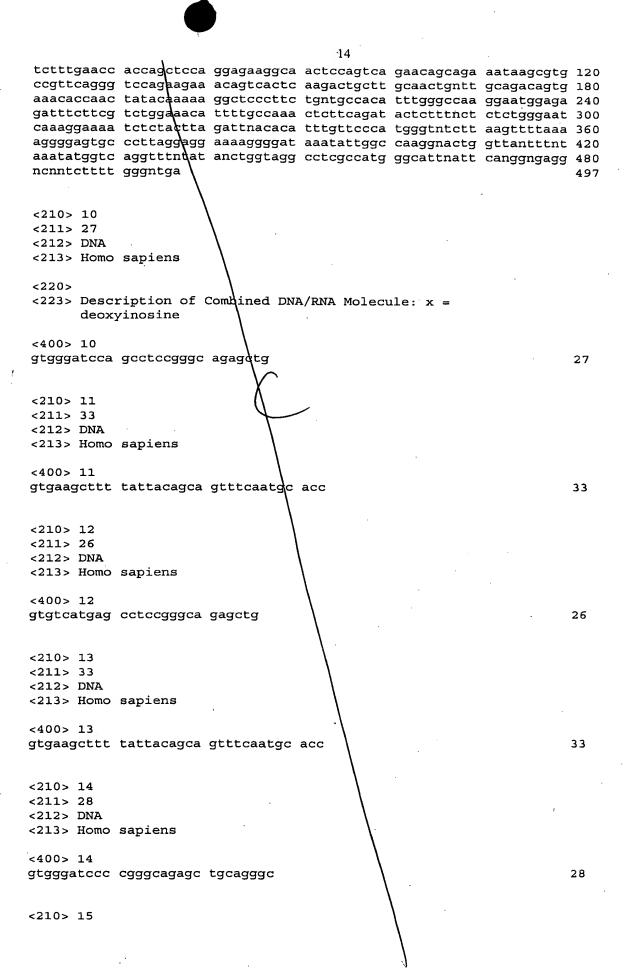
```
<222> (300)
<223> n equals a, t \setminus g, or c
<220>
<221> misc_feature
<222> (320)
<223> n equals a, t, g or c
<220>
<221> misc_feature
<222> (335)
<223> n equals a, t, g, dr c
<400> 7
ggntaactet cetgaggggt gagecaagee etgecatgta gtgeacgeag gacateanea 60
aacacannnn ncaggaaata atccattccc tgtggtcact tattctaaag gccccaacct 120
tcaaagttca agtagtgata tggatgactc cacagaaagg gagcagtcac gccttacttc 180
ttgccttaag aaaagagaag aaatgaaact gnaaggagtg tgtttccatc ctcccacgga 240
aggaaagccc ctctntccga tcctccaaag acggaaagct gctggctgca accttgntgn 300
tggcattgtg ttcttgctgn ctcaaggtgg tgttntt
<210> 8
<211> 509
<212> DNA
<213> Homo sapiens
<220>
<223> Description of Combined DNAVRNA Molecule: n equals
      a, t, g, or c
<220>
<221> misc feature
<222> (10)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (13)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (209)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (315)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (322)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (325)
<223> n equals a, t, g, or c
```

```
<220>
<221> misc feature
<222> (334)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (343)
<223> n equals a, t, g
                        or c
<220>
<221> misc feature
<222> (347)
<223> n equals a, t, g, dr c
<220>
<221> misc_feature
<222> (351)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (356)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (409) .. (410)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (416)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (422)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (424)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (426)..(427)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (429)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222'> (431)
<223> n equals a, t, g, or c
```

```
<220>
<221> misc feature
<222> (433)
<223> n equals a,t, g, or c
<220>
<221> misc feature
<222> (438)..(439)
<223> n equals a, t,\g, or c
<220>
<221> misc_feature
<222> (443)..(444)
<223> n equals a, t, g,\or c
<220>
<221> misc_feature
<222> (446)..(447)
<223> n equals a, t, g, or\c
<220>
<221> misc_feature
<222> (449)..(450)
<223> n equals a, t, g, or (c
<220>
<221> misc_feature
<222> (452)..(453)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (458)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (461)..(462)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (466)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (469)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (471)..(472)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (474)
<223> n equals a, t, g, or c
```

```
<220>
<221> misc feature
<222> (478)..(481)
\langle 223 \rangle n equals a, t
                     g, or c
<220>
<221> misc feature
<222> (496)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (498)
<223> n equals a, t, g,
<220>
<221> misc_feature
<222> (504)
<223> n equals a, t, g, or
<400> 8
aattoggoan agnaaactgg tta¢tt\tt atatatggto aggttttata tactgataag 60
acctacgcca tgggacatct agttcagagg aagaaggtcc atgtctttgg ggatgaattg 120
agtetggtga etttgttteg atgbattdaa aatatgeetg aaacaetaee caataattee 180
tgctattcag ctggcattgc aaaactggha ggaaggagat gaactccaac ttgcaatacc 240
aggggaaaat gcacaattat cactgggadg gagatgttca cattttttgg gtgccattga 300
aactgctgtg acctnettae ancangtget gttngctatt ttnectneet nttetntggt 360
aacctcttag gaaggaagga ttcttaactg ggaaataacc caaaaaaann ttaaangggt 420
angngnnana ngnggggnng ttnncnngnn\gnnttttngg nntatnttnt nntngggnnn 480
ngtaaaaatg gggccnangg gggnttttt
<210> 9
<211> 497
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (168)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (213)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (288)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (325)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
```

```
<222> (346)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (406)
<223> n equals a, t,\g, or c
<220>
<221> misc feature
<222> (415)
<223> n equals a, t, g \setminus or c
<220>
<221> misc feature
<222> (419)
<223> n equals a, t, g, of c
<220> .
<221> misc_feature
<222> (437)
<223> n equals a, t, g, or d
<220>
<221> misc feature
<222> (442)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (467)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (473)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (476)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (481)
<223> n equals a, t, g, or c
<220>
<221> misc_feature
<222> (483)..(484)
<223> n equals a, t, g, or c
<220>
<221> misc feature
<222> (494)
<223> n equals a, t, g, or c
<400> 9
aatteggeae gageaaggee ggeetggagg aageteeage/tgteacegeg ggaetgaaaa 60
```



| | | | | | | | | | 13 | | | | |
|--------------|------------------------------|------------|----------------|------|------|------|------------------|-----------|----|--|--|------------------|-----|
| <21 | 1 > 3 2 > D 3 > H | NA | \ sapi | ens | | | | | | | | | |
| | 0> 1 ggat | | tatt | adag | ca g | tttc | aatg | c ac | c | | | | 33 |
| <21 <21 | 0> 1 1> 1 2> D 3> H | 29 NA | sapi | ens | \ | | | | | | | | |
| gcgg | | ccg tgc | | | | | | | | | | ttetee ggggae | |
| <213 | 0> 1' 1> 3(2> Di | 0 NA | | | | | | . 1. • | | | | | |
| <400 | 0> 1' | 7 | sapi taca | | tt t | caat | gcaco | | | | | | 30 |
| <211 <212 | 0> 18 L> 90 2> Di | EC AVA | | | | | | · · · | : | | | | |
| <220 <221 |)> L> CI | os | sapie (798) | | | | | | | | | • | |
| atg | | gac | | | | | gag Glu | | | | | | 48 |
| | | | | | | | ctg Leu | | | | | | 96 |
| | | | | | | | cga Arg 40 | | | | | | 144 |
| | | | | | | | ctg Leu | | | | | | 192 |
| | | | | | | | ctg Leu | | | | | | 240 |
| | | | | | | | gcg Ala | | | | | | 288 |

| | \ 85 | | 16 90 | 95 | |
|--|-----------------------------------|-----------------------------------|-----------------------------------|---|--------|
| | | | | acc gcg gga ctg Thr Ala Gly Leu 110 | |
| aaa atc ttt Lys Ile Phe 115 | gaa cca cca Glu Pro Pro | gct cca gga Ala Pro Gly 120 | gaa ggc aac Glu Gly Asn | tcc agt cag aac Ser Ser Gln Asn 125 | 384 |
| | | | | aca gga tct tac Thr Gly Ser Tyr | 432 |
| | | | | agt gcc cta gaa Ser Ala Leu Glu 160 | 480 |
| | | | | tac ttt ttt ata Tyr Phe Phe Ile 175 | 528 |
| | | | Thr Tyr Ala | atg gga cat cta Met Gly His Leu 190 | 576 |
| | | | Gly Asp Glu | ttg agt ctg gtg Leu Ser Leu Val 205 | 624 |
| | Arg Cys Ile | | | cta ccc aat aat Leu Pro Asn Asn | 672 |
| | | | | gga gat gaa ctc Gly Asp Glu Leu 240 | 720 |
| caa ctt gca Gln Leu Ala | ata cca aga Ile Pro Arg 245 | gaa aat gca Glu Asn Ala | caa ata tca Gln tle Ser 250 | ctg gat gga gat Leu Asp Gly Asp 255 | 768 |
| Val Thr Phe | | | | tt acaccatgtc | 818 |
| tgtagctatt t | tcctccctt tc | tctgtacc tct | aagaaga \aaga | atctaa ctgaaaata | ac 878 |
| caaaaaaaaa a | aaaaaaaaa aa | aaa | | | 903 |
| <210> 19 <211> 266 <212> PRT <213> Homo s | apiens | | | | |
| <400> 19 Met Asp Asp 1 | Ser Thr Glu . | Arg Glu Gln | Ser Arg Leu | thr Ser Cys Leu 15 | • |

Lys Lys Arg Glu Glu Met Lys Leu Lys Glu Cys Val Ser Ile Leu Pro 20 25 30

Arg Lys Glu Ser Pro Ser Val Arg Ser Ser Lys Asp Gly Lys Leu Leu
35 40 45

Ala Ala Thr Leu Leu Leu Ala Leu Leu Ser Cys Cys Leu Thr Val Val
50 60

Ser Phe Tyr Gln Val Ala Leu Gln Gly Asp Leu Ala Ser Leu Arg
65 70 75 80

Ala Glu Leu Gln Gly His Nis Ala Glu Lys Leu Pro Ala Gly Ala Gly
85 90 95

Ala Pro Lys Ala Gly Leu Gly Glu Ala Pro Ala Val Thr Ala Gly Leu
100 105 110

Lys Ile Phe Glu Pro Pro Ala Pro Gly Glu Gly Asn Ser Ser Gln Asn
115 120 125

Ser Arg Asn Lys Arg Ala Val the Gly Pro Glu Glu The Gly Ser Tyr

Thr Phe Val Pro Trp Leu Leu Ser\Phe Lys Arg Gly Ser Ala Leu Glu
145 150 155 160

Glu Lys Glu Asn Lys Ile Leu Val Lys Glu Thr Gly Tyr Phe Phe Ile 165 170 175

Tyr Gly Gln Val Leu Tyr Thr Asp Lys Thr Tyr Ala Met Gly His Leu 180 185 190

Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu Ser Leu Val 195 200 205

Thr Leu Phe Arg Cys Ile Gln Asn Met Pro Glu Thr Leu Pro Asn Asn 210 215 220

Ser Cys Tyr Ser Ala Gly Ile Ala Lys Leu Glu Glu Gly Asp Glu Leu 225 230 240

Gln Leu Ala Ile Pro Arg Glu Asn Ala Gln tle Ser Leu Asp Gly Asp
245 250 255

Val Thr Phe Phe Gly Ala Leu Lys Leu Leu 260 265

<210> 20

<211> 136

<212> PRT

<213> Homo sapiens

<400> 20

His Ser Val Leu His Leu Val Pro Ile Asn Ala Thr Ser Lys Asp Asp

1 5 10 15

Ser Asp Val Thr Glu Val Met Trp Gln Pro Ala Leu\Arg Arg Gly Arg

Gly Leu Gln Ala Gln Gly Tyr Gly Val Arg Ile Gln Asp Ala Gly Val 40

25

Tyr Leu Leu Tyr Ser Gln Val Leu Phe Gln Asp Val Thr Phe Thr Met

Gly Gln Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu Thr Leu Phe

Arg Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala Tyr Asn Ser

Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp Ile Leu Ser

Val Ile Ile Pro Arg Ala Arg \Ala Lys Leu Asn Leu Ser Pro His Gly 120

Thr Phe Leu Gly Phe Val Lys L

atggctgttc agggtccgga agaaaccgtt actcaggact gccttcagct gatcgcagac 60 totgaaacto ogacoatoca gaaaggttot tadacetttg ttoottggot gotttottto 120 aaacgtggtt ctgccctgga agagaaagaa aackaaatcc tggttaaaga aactggttac 180 ttetttatet aeggteaggt tetttaeaet gataagaeet aegeeatggg teaeetgatt 240 cagegtaaga aagtteaegt ttteggtgae gagetgtete tggttaetet gtttegetge 300 atteagaaca tgeeggaaae tetteetaae aaet¢etget aetetgetgg categeaaaa 360 ctggaagagg gtgatgaact gcagctggca attcctcgtg aaaacgcaca aatttctctg 420 gacggtgatg taaccttctt tggtgcactg aaacttctgt aa

<210> 22

<211> 1040

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (1)..(468)

ege gtg gta gae ete tea get eet eet gea eea tge etg eet gga tge 48 Arg Val Val Asp Leu Ser Ala Pro Pro Ala Pro dys Leu Pro Gly Cys

cgc cat tot caa cat gat gat aat gga atg aac ott aga aac aga act 96 Arg His Ser Gln His Asp Asp Asn Gly Met Asn Led Arg Asn Arg Thr

tac aca ttt gtt cca tgg ctt ctc agc ttt aaa aga gga aat gcc ttg Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys Arg dly Asn Ala Leu

| ųΩ | |
|----|--|
| ĻΠ | |
| ĊŌ | |
| ā | |
| īŪ | |
| 10 | |
| Ü | |
| 15 | |
| 1 | |
| m | |
| I | |
| Ö | |
| | |
| | |
| | |
| | |

| | | | | \ | | | | | | | | 13 | | | | |
|------------------|------------|-------------------|-------------------|------------------|------------------|------------|-------------------|-------------------|------------------|------------------|------------|-------------------|-------------------|------------------|------------------|------|
| | | | | | | | | | | | | | | ttc Phe | | 192 |
| atc Ile 65 | tac Tyr | agc Ser | cag Gln | gtt Val | cta Leu 70 | tac Tyr | acg Thr | gac Asp | ccc Pro | atc Ile 75 | ttt Phe | gct Ala | atg Met | ggt Gly | cat His 80 | 240 |
| gtc Val | atc Ile | cag Gln | agg Arg | aag Lys 85 | aaa Lys | yta Val | cac His | gtc Val | ttt Phe 90 | gly aaa | gac Asp | gag Glu | ctg Leu | agc Ser 95 | ctg Leu | 288 |
| gtg Val | acc Thr | ctg Leu | ttc Phe 100 | cga Arg | tgt Cys | att | cag Gln | aat Asn 105 | atg Met | ccc Pro | aaa Lys | aca Thr | ctg Leu 110 | ccc Pro | aac Asn | 336 |
| aat Asn | tcc Ser | tgc Cys 115 | tac Tyr | tcg Ser | gct Ala | ggc Gly | atc Ile 130 | gcg Ala | agg Arg | ctg Leu | gaa Glu | gaa Glu 125 | gga Gly | gat Asp | gag Glu | 384 |
| | | | | | | | | | | | | | | aac Asn | | 432 |
| | | | | | | | | | ctg Leu | | taa | ctca | cttg | jct | | 478 |
| ggag | tgcg | ıtg a | taca | ctto | c ct | cgtc | ttct | : dtg | gtacc | tcc | gagg | gaga | aa c | cagac | gactg | 538 |
| gaaa | aact | aa a | agat | 9999 | ra aa | ıgccg | rtcag | 1 c3/6 | aagt | ttt | ctcg | rtgac | :cc 9 | ıttga | atctg | 598 |
| atco | aaac | ca g | gaaa | tata | a ca | gaca | gcca | caa | cga | agt | gtgo | catg | tg a | ıgtta | tgaga | 658 |
| aacg | gago | cc g | cgct | caga | a ag | accg | gatg | agg | Jaagga | ccg | tttt | ctcc | ag t | cctt | tgcca | 718 |
| acac | gcac | cg c | aacc | ttgo | t tt | ttgc | cttg | ggt | ga¢a | cat | gtto | agaa | tg c | aggg | agatt | 778 |
| taat | tgtt | tt g | cgat | ttgc | c at | gaga | .agag | ggc | cca | aac | tgca | ggtc | ac t | gaag | cattc | 838 |
| acgo | taag | rtc t | cagg | atțt | a ct | ctcc | atta | tca | tgct | aag | taca | .caca | .cg c | tctt | ttcca | 898 |
| ggta | atac | ta t | ggga | tact | a tg | gaaa | ggtt | gtt | tgtt | det | aaat | ctag | aa g | tett | gaact | 958 |
| ggca | atag | ac a | aaaa | tcct | t at | aaat | tcaa | gtg | rtaaa | ata | aact | taat | ta a | aaag | gttta | 1018 |
| agtg | tgaa | aa a | aaaa | aaaa | a aa | • | | | | | - | | | | | 1040 |
| | | | | | | | | | | | | | | | | |

<210> 23

<211> 155

<212> PRT

<213> Homo sapiens

<400> 23

Arg Val Val Asp Leu Ser Ala Pro Pro Ala Pro Cys Leu Pro Gly Cys

| 1 10 15 | |
|---|-------------------------|
| Arg His Ser Gln His Asp Asp Asn Gly Met Asn Leu Arg Asn Arg Thr | |
| 20 \ 25 30 | |
| Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys Arg Gly Asn Ala Leu | |
| 35 \ 40 45 | |
| Glu Glu Lys Glu Asn Lys Ile Val Val Arg Gln Thr Gly Tyr Phe Phe | |
| 50 \ 55 60 | |
| Ile Tyr Ser Gln Val Leu Tyr Thr Asp Pro Ile Phe Ala Met Gly His | |
| | |
| · - | |
| Val Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu Ser Leu | |
| 85 \ 90 95 | |
| Val Thr Leu Phe Arg Cyp Ile Gln Asn Met Pro Lys Thr Leu Pro Asn | |
| 100 \ 105 \ 110 | |
| Asn Ser Cys Tyr Ser Ala Gly Ile Ala Arg Leu Glu Glu Gly Asp Glu | |
| 115 \ 120 125 | |
| Ile Gln Leu Ala Ile Pro Arg Glu Asn Ala Gln Ile Ser Arg Asn Gly | |
| 130 135 140 | |
| Asp Asp Thr Phe Phe Gly Ala Leu Lys Leu Leu | |
| | |
| 145 150 \ 155 | |
| \ | |
| | |
| √ \ . | |
| <210> 24 | |
| <211> 26 | |
| <212> DNA / \ | |
| <213> Homo sapiens (\ | |
| + | |
| <400> 24 | |
| | 26 |
| ccaccagete caggagaagg caacte \ | 26 |
| | • |
| | |
| <210> 25 | |
| <211> 19 \ | |
| <212> DNA \ | |
| <213> Homo sapiens | |
| \ | |
| <400> 25 | |
| accgcgggac tgaaaatct \ | 19 |
| \ | |
| | |
| 121.0. 26 | |
| <210> 26 | |
| <211> 23 | |
| <212> DNA \ | |
| <213> Homo sapiens | |
| \ | |
| | |
| <400> 26 | |
| <pre><400> 26 cacgcttatt tctgctgttc tga</pre> | 23 |
| | 23 |
| | 23 |
| cacgcttatt tctgctgttc tga | 23 |
| <210> 27 | 23 |
| <210> 27 <211> 657 | 23 |
| <210> 27 <211> 657 <212> DNA | 23 |
| <210> 27 <211> 657 | 23 |
| <210> 27 <211> 657 <212> DNA | 23 |
| <210> 27 <211> 657 <212> DNA | 23 |
| <pre>cacgcttatt tctgctgttc tga <210> 27 <211> 657 <212> DNA <213> Homo sapiens <400> 27</pre> | |
| <pre>cacgcttatt tctgctgttc tga <210> 27 <211> 657 <212> DNA <213> Homo sapiens <400> 27 taccaggtgg cggccgtgca aggggacctg gccagcctcc gggcagagct gcagggccac</pre> | 60 |
| <pre>cacgcttatt tctgctgttc tga <210> 27 <211> 657 <212> DNA <213> Homo sapiens <400> 27 taccaggtgg cggccgtgca aggggacctg gccagcctcc gggcagagct gcagggccaccacgcggaga agctgccagc aagagcaaga gccccaagg cdggtctggg ggaagctcca</pre> | 60 120 |
| <pre>cacgcttatt tctgctgttc tga <210> 27 <211> 657 <212> DNA <213> Homo sapiens <400> 27 taccaggtgg cggccgtgca aggggacctg gccagcctcc gggcagagct gcagggccaccagggaga agctgcagc aagagcaaga gccccaagg cdggtctggg ggaagctccagctgtcaccg caggactga aatctttgaa ccaccagct caggagaagg caactccagt</pre> | 60 120 180 |
| <pre>cacgcttatt tctgctgttc tga <210> 27 <211> 657 <212> DNA <213> Homo sapiens <400> 27 taccaggtgg cggccgtgca aggggacctg gccagcctcc gggcagagct gcagggccaccacgcggaga agctgccagc aagagcaaga gccccaagg cdggtctggg ggaagctcca</pre> | 60 120 180 240 |

ccatggcttc tcagcttta aaagggaagt gccctagaag aaaaagagaa taaaatattg 360 gtcaaagaaa ctggttactt ttttatatat ggtcaggttt tatacactga taagacctat 420 gccatgggac atctaattca gaggaaaaaa gtccatgtct ttggggatga attgagtctg 480 gtgactttgt ttcgatgtat tcaaaatatg cctgaaacac tacccaataa ttcctgctat 540 tcagctggca ttgcaaaact ggaagaagga gatgaacttc aacttgcaat accacgagaa 600 aatgcacaaa tatcactgga tggagatgtc acattttttg gtgccctcaa actgctg 657

<210> 28

<211> 219

<212> PRT

<213> Homo sapiens

<400> 28

Tyr Gln Val Ala Ala Val Gln Gly Asp Leu Ala Ser Leu Arg Ala Glu

1 1 15

Leu Gln Gly His His Ala Glu Nys Leu Pro Ala Arg Ala Arg Ala Pro
20 25 30

Lys Ala Gly Leu Gly Glu Ala Pro Ala Val Thr Ala Gly Leu Lys Ile
35 45

Phe Glu Pro Pro Ala Pro Gly Glu Gly Asn Ser Ser Gln Ser Ser Arg
50 55 60

Asn Lys Arg Ala Ile Gln Gly Ala Glu Glu Thr Val Ile Gln Asp Cys
65 70 75 80

Leu Gln Leu Ile Ala Asp Ser Glu Thr\Pro Thr Ile Gln Lys Gly Ser
85 95

Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys Arg Gly Ser Ala Leu 100 105 110

Glu Glu Lys Glu Asn Lys Ile Leu Val Lys Glu Thr Gly Tyr Phe Phe 115 120 125

Ile Tyr Gly Gln Val Leu Tyr Thr Asp Lys Thr Tyr Ala Met Gly His
130 135 140

Leu Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu Ser Leu 145 150 155 160

Val Thr Leu Phe Arg Cys Ile Gln Asn Met Pro Glu Thr Leu Pro Asn 165 170 175

Asn Ser Cys Tyr Ser Ala Gly Ile Ala Lys Leu Glu Glu Gly Asp Glu
180 185 190

Leu Gln Leu Ala Ile Pro Arg Glu Asn Ala Gln Ile Ser Leu Asp Gly
195 200 205

Asp Val Thr Phe Phe Gly Ala Leu Lys Leu Leu 210 215

<210> 29

<211> 657

<212> DNA

<213> Homo sapiens

<400> 29
taccaggtgg cggcgtgca aggggacctg gccagcctcc gggcagagct gcagagccac 60
cacgcggaga agctgcagc aagagcaaga gccccaaagg ccggtctggg ggaagctcca 120
gctgtcaccg cgggactgaa aatctttgaa ccaccagctc caggagaagg caactccagt 180
cagagcagca gaaataagcg tgctattcag ggtgcagaag aaacagtcat tcaagactgc 240
ttgcaactga ttgcagatag tgaaacacca actatacaaa aaggatctta cacatttgtt 300
ccatggcttc tcagctttaa aaggggaagt gccctagaag aaaaagagaa taaaatattg 360
gtcaaagaaa ctggttactt ttttatatat ggtcaggttt tatacactga taagacctat 420
gccatgggac atctaattca gaggaaaaaa gtccatgtct ttggggatga attgagtctg 480
gtgactttgt ttcgatgtat tcaaaatatg cctgaaacac tacccaataa ttcctgctat 540
tcagctggca ttgcaaaact gggaagaggg gatgaacttc aacttgcaat accacgagaa 600
aatgcacaaa tatcactgga tggagatgtc acattttttg gtgccctcaa actgctg 657

<400> 30

<213> Homo sapiens

Tyr Gln Val Ala Ala Val Gln Gly Asp Leu Ala Ser Leu Arg Ala Glu
1 10 15

Leu Gln Ser His His Ala Glu Lys Leu Pro Ala Arg Ala Arg Ala Pro
20 25 30

Lys Ala Gly Leu Gly Glu Ala Pro Ala Val Thr Ala Gly Leu Lys Ile 35 40 45

Phe Glu Pro Pro Ala Pro Gly Glu Gly Asn Ser Ser Gln Ser Ser Arg
50 55 60

Asn Lys Arg Ala Ile Gln Gly Ala Glu Glu Thr Val Ile Gln Asp Cys
65 70 75 80

Leu Gln Leu Ile Ala Asp Ser Glu Thr\Pro Thr Ile Gln Lys Gly Ser

Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys Arg Gly Ser Ala Leu 100 105 110

Glu Glu Lys Glu Asn Lys Ile Leu Val Lys Glu Thr Gly Tyr Phe Phe
115 120 125

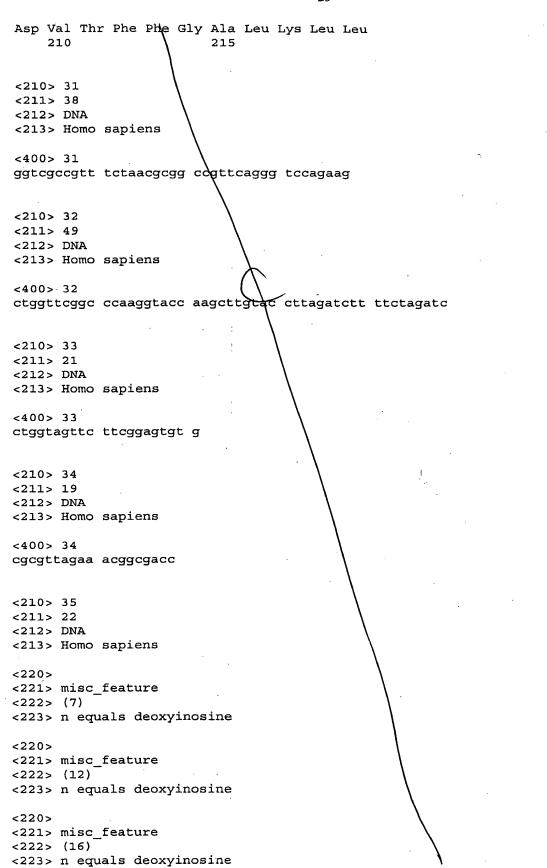
Ile Tyr Gly Gln Val Leu Tyr Thr Asp Lys Thr Tyr Ala Met Gly His

Leu Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu Ser Leu 145 150 155 160

Val Thr Leu Phe Arg Cys Ile Gln Asn Met Pro Glu Thr Leu Pro Asn
165 170 175

Asn Ser Cys Tyr Ser Ala Gly Ile Ala Lys Leu Glu Glu Gly Asp Glu
180 185 190

Leu Gln Leu Ala Ile Pro Arg Glu Asn Ala Gln Ile Ser Leu Asp Gly
195 200 205



```
<400> 35
taccagntgg cngccntgca ag
                                                                    22
<210> 36
<211> 22
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (3)
<223> n equals deoxyinosine
<220>
<221> misc feature
<222> (14)
<223> n equals deoxyinosine
<220>
<221> misc feature
<222> (16)..(17)
<223> n equals deoxyinosine
<400> 36
gtnacagcag tttnanngca cc
                                                                   22
<210> 37
<211> 866
<212> DNA
<213> Mus musculus
<400> 37
atggatgagt ctgcaaagac cctgccacca ccgtg&ctct gtttttgctc cgagaaagga 60
gaagatatga aagtgggata tgatcccatc actccgcaga aggaggaggg tgcctggttt 120
gggatetgea gggatggaag getgetgget getacedtee tgetggeeet gttgteeage 180
agtttcacag cgatgtcctt gtaccagttg gctgccttgc.aagcagacct gatgaacctg 240
cgcatggagc tgcagagcta ccgaggttca gcaacacdag ccgccgcggg tgctccagag 300
ttgaccgetg gagtcaaact cetgacaccg gcagetecte gaccecacaa etccagcege 360
ggccacagga acagacgege ettecaggga ccagaggada cagaacaaga tgtagacete 420
tcageteete etgeaceatg eetgeetgga tgeegeeatt eteaacatga tgataatgga 480
atgaacctca gaaacatcat tcaagactgt ctgcagctga ttgcagacag cgacacgccg 540
gccttggagg agaaagagaa caaaatagtg gtgaggcaaa \caggctattt cttcatctac 600
agccaggttc tatacacgga ccccatcttt gctatgggtc atgtcatcca gaggaagaaa 660
gtacacgtet ttggggacga getgageetg gtgaceetgt tcggatgtat teagaatatg 720
cccaaaacac tgcccaacaa ttcctgctac tcggctggca tdgcgaggct ggaagaagga 780
gatgagatte agettgeaat teetegggag aatgeacaga tt/caegeaa eggagaegae 840
accttctttg gtgccctaaa actgct
<210> 38
<211> 177
<212> PRT
<213> Mus musculus
<400> 38
Met Asp Ser Ala Lys Thr Cys Cys Cys Ser Lys Gly Asp Met Lys Val
```





Gly Tyr Asp Thr Lys Gly Ala Trp Gly Cys Arg Asp Gly Arg Ala Ala 20 \downarrow 25 30

Thr Ala Ser Ser Ser Thr Ala Met Ser Tyr Ala Ala Ala Asp Met Asn 35 40 45

Arg Met Ser Tyr Arg Gly Ser Ala Thr Ala Ala Ala Gly Ala Thr Ala
50 60

Gly Val Lys Thr Ala Ala Arg His Asn Ser Ser Arg Gly His Arg Asn 65 70 75 80

Arg Arg Ala Gly Thr Asp Val Asp Ser Ala Ala Cys Gly Cys Arg His
85
90
95

Ser His Asp Asp Asn Gly Met Asn\Arg Asn Asp Cys Ala Asp Ser Asp
100 110

Thr Ala Lys Asn Lys Val Val Arg Thr Gly Tyr Tyr Ser Val Tyr Thr
115 120 125

Asp Ala Met Gly His Val Arg Lys Lys Val His Val Gly Asp Ser Val
130 135 140

Thr Arg Cys Asn Met Lys Thr Asn Asn Ser Cys Tyr Ser Ala Gly Ala
145 150 155 160

Arg Gly Asp Ala Arg Asn Ala Ser Arg Ash Gly Asp Asp Thr Gly Ala
165 170 175

Lys